RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _____ 09/981, 151 D

Source: _____ IFW16

Date Processed by STIC: _____ 08/22/2005

ENTERED

IFW16

RAW SEQUENCE LISTING   DATE: 08/22/2005
PATENT APPLICATION: US/09/981,151D   TIME: 10:02:32

Input Set : **D:\Cura 468 SEQ list 0705.txt**
Output Set: **N:\CRF4\08222005\I981151D.raw**

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 3  <110> APPLICANT: Guo, Xiaojia
 5  <120> TITLE OF INVENTION: Proteins and Nucleic Acids Encoding Same
 7  <130> FILE REFERENCE: 21402-168
 9  <140> CURRENT APPLICATION NUMBER: 09/981,151D
10  <141> CURRENT FILING DATE: 2001-10-16
12  <150> PRIOR APPLICATION NUMBER: 60/241,040
13  <151> PRIOR FILING DATE: 2000-10-17
15  <150> PRIOR APPLICATION NUMBER: 60/241,058
16  <151> PRIOR FILING DATE: 2000-10-17
18  <150> PRIOR APPLICATION NUMBER: 60/241,063
19  <151> PRIOR FILING DATE: 2000-10-17
21  <150> PRIOR APPLICATION NUMBER: 60/241,243
22  <151> PRIOR FILING DATE: 2000-10-17
24  <150> PRIOR APPLICATION NUMBER: 60/242,152
25  <151> PRIOR FILING DATE: 2000-10-20
27  <150> PRIOR APPLICATION NUMBER: 60/242,482
28  <151> PRIOR FILING DATE: 2000-10-23
30  <150> PRIOR APPLICATION NUMBER: 60/242,611
31  <151> PRIOR FILING DATE: 2000-10-23
33  <150> PRIOR APPLICATION NUMBER: 60/242,612
34  <151> PRIOR FILING DATE: 2000-10-23
36  <150> PRIOR APPLICATION NUMBER: 60/242,880
37  <151> PRIOR FILING DATE: 2000-10-24
39  <150> PRIOR APPLICATION NUMBER: 60/242,881
40  <151> PRIOR FILING DATE: 2000-10-24
42  <150> PRIOR APPLICATION NUMBER: 60/259,028
43  <151> PRIOR FILING DATE: 2000-12-29
45  <150> PRIOR APPLICATION NUMBER: 60/269,813
46  <151> PRIOR FILING DATE: 2001-02-20
48  <150> PRIOR APPLICATION NUMBER: 60/286,324
49  <151> PRIOR FILING DATE: 2001-04-25
51  <150> PRIOR APPLICATION NUMBER: 60/294,108
52  <151> PRIOR FILING DATE: 2001-05-29
54  <150> PRIOR APPLICATION NUMBER: 60/303,968
55  <151> PRIOR FILING DATE: 2001-07-09
57  <160> NUMBER OF SEQ ID NOS: 166
59  <170> SOFTWARE: PatentIn Ver. 2.1
61  <210> SEQ ID NO: 1
62  <211> LENGTH: 2997
63  <212> TYPE: DNA
64  <213> ORGANISM: Homo sapiens
66  <220> FEATURE:
67  <221> NAME/KEY: misc_feature
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(pg.6)

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 84 tccggacagc tggaggcgag tcccccgcgg ctcctctccc gcggaccccg ccgtctcacc 180
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 87 aaagggcggg acatggatga agctggaaac catcgttctc agcaaactaa cacaggaaca 360
 88 gaaaaccaaa cactgcatgt tctcactcaa tatgacctgg tctctgccta cgaggttgac 420
 89 cacaggggcg attacgtgtc ccatgaaatc atgcaccatc agcggcggag aagagcagtg 480
 90 gccgtgtccg aggttgagtc tcttcacctt cggctgaaag gccccaggca cgacttccac 540
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118 aaatgtcccc gggacagtgt tgacttccgt gctgctcagt gtgccgagca caacagcaga 2220
119 cgattcagag ggcggcacta caagtggaag cctcaggact tatgcaaact ctactgtatc 2280
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RAW SEQUENCE LISTING   DATE: 08/22/2005
PATENT APPLICATION: US/09/981,151D   TIME: 10:02:32

Input Set : D:\Cura 468 SEQ list 0705.txt
Output Set: N:\CRF4\08222005\I981151D.raw

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122 cttggatctg atgctgttga agacgtctgt ggggtgtgta acgggaataa ctcagcctgc 2460
123 acgattcaca ggggtctcta.caccaagcac caccacacca accattatca catggtcacc 2520
124 attccttctg gagcccggag tatccgcatc tatgaaatga acgtctctac ctcctacatt 2580
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126 ggccggtaca aattttcggg cactactttc gactacagac ggtcctataa tgagcccgag 2700
127 aacttaatcg ctactggacc aaccaacgag acactgattg tggagctgct gtttcaggga 2760
128 aggaacccgg gtgttgcctg ggaatactcc atgcctcgct gggggaccga gaagcagccc 2820
129 cctgcccagc ccagctacac ttgggccatc gtgcgctctg agtgctccgt gtcctgcgga 2880
130 gggggtaggt gccttccagt gctgctcctg gaggcagcat gtcagccttc agccactgcg 2940
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135 <211> LENGTH: 986
136 <212> TYPE: PRT
137 <213> ORGANISM: Homo sapiens
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140 <221> NAME/KEY: VARIANT
141 <222> LOCATION: (283)
142 <223> OTHER INFORMATION: Wherein Xaa is any amino acid.
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145 <221> NAME/KEY: VARIANT
146 <222> LOCATION: (792)
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154              20                  25                  30
156 Arg Gly Asn Arg Gly Ser Gly Gln Leu Glu Ala Ser Pro Pro Arg Leu
157          35                  40                  45
159 Leu Ser Arg Gly Pro Arg Arg Leu Thr Ala Met Ser Pro Leu Phe Ser
160      50                  55                  60
162 Ala Gly Thr Cys Val Arg His Gly Thr Arg Ser Gly Ser Ala Trp Glu
163 65                  70                  75                  80
165 Pro Glu Arg Pro Ala Ser Ser Ser Thr Arg Gly Ala Ala Gly Leu Asp
166                  85                  90                  95
168 Gly Lys Gly Arg Asp Met Asp Glu Ala Gly Asn His Arg Ser Gln Gln
169              100                 105                 110
171 Thr Asn Thr Gly Thr Glu Asn Gln Thr Leu His Val Leu Thr Gln Tyr
172          115                 120                 125
174 Asp Leu Val Ser Ala Tyr Glu Val Asp His Arg Gly Asp Tyr Val Ser
175          130                 135                 140
177 His Glu Ile Met His His Gln Arg Arg Arg Ala Val Ala Val Ser
178 145                 150                 155                 160
180 Glu Val Glu Ser Leu His Leu Arg Leu Lys Gly Pro Arg His Asp Phe
181                  165                 170                 175
183 His Met Asp Leu Arg Thr Ser Ser Ser Leu Val Ala Pro Gly Phe Ile
184              180                 185                 190
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        186 Val Gln Thr Leu Gly Lys Thr Gly Thr Lys Ser Val Gln Thr Leu Pro
        187         195             200             205
        189 Pro Glu Asp Phe Cys Phe Tyr Gln Gly Ser Leu Arg Ser His Arg Asn
        190     210             215             220
        192 Ser Pro Ser His Gly Gly Lys Phe Cys Glu Gly Ser Thr Arg Thr Leu
        193 225             230             235                 240
        195 Lys Leu Cys Asn Ser Gln Lys Cys Pro Arg Asp Ser Val Asp Phe Arg
        196             245             250                 255
        198 Ala Ala Gln Cys Ala Glu His Asn Ser Arg Arg Phe Arg Gly Arg His
        199             260             265             270
W-->    201 Tyr Lys Trp Lys Pro Tyr Thr Gln Val Glu Xaa Asp Leu Cys Lys Leu
        202         275             280             285
        204 Tyr Cys Ile Ala Glu Gly Phe Asp Phe Phe Phe Ser Leu Ser Asn Lys
        205         290             295             300
        207 Val Lys Asp Gly Thr Pro Cys Ser Glu Asp Ser Arg Asn Val Cys Ile
        208 305             310             315                 320
        210 Asp Gly Ile Cys Glu Leu Ser Val Val Ser Thr Ser Ala His Met Pro
        211             325             330             335
        213 Gln Pro Pro Lys Glu Asp Leu Phe Ile Leu Pro Asp Glu Tyr Lys Ser
        214             340             345             350
        216 Cys Leu Arg His Lys Arg Ser Leu Leu Arg Ser His Arg Asn Glu Glu
        217         355             360             365
        219 Leu Asn Val Glu Thr Leu Val Val Val Asp Lys Lys Met Met Gln Asn
        220     370             375             380
        222 His Gly His Glu Asn Ile Thr Thr Tyr Val Leu Thr Ile Leu Asn Met
        223 385             390             395                 400
        225 Val Ser Ala Leu Phe Lys Asp Gly Thr Ile Gly Gly Asn Ile Asn Ile
        226             405             410                 415
        228 Ala Ile Val Gly Leu Ile Leu Leu Glu Asp Glu Gln Pro Gly Leu Val
        229             420             425             430
        231 Ile Ser His His Ala Asp His Thr Leu Ser Ser Phe Cys Gln Trp Gln
        232         435             440             445
        234 Ser Gly Leu Met Gly Lys Asp Gly Thr Arg His Asp His Ala Ile Leu
        235     450             455             460
        237 Leu Thr Gly Leu Asp Ile Cys Ser Trp Lys Asn Glu Pro Cys Asp Thr
        238 465             470             475                 480
        240 Leu Gly Phe Ala Pro Ile Ser Gly Met Cys Ser Lys Tyr Arg Ser Cys
        241             485             490             495
        243 Thr Ile Asn Glu Asp Thr Gly Leu Gly Leu Ala Phe Thr Ile Ala His
        244             500             505             510
        246 Glu Ser Gly His Asn Phe Gly Met Ile His Asp Gly Glu Gly Asn Met
        247         515             520             525
        249 Cys Lys Lys Ser Glu Gly Asn Ile Met Ser Pro Thr Leu Ala Gly Arg
        250         530             535             540
        252 Asn Gly Val Phe Ser Trp Ser Pro Cys Ser Arg Gln Tyr Leu His Lys
        253 545             550             555                 560
        255 Phe Leu Ser Thr Ala Gln Ala Ile Cys Leu Ala Asp Gln Pro Lys Pro
        256             565             570             575
        258 Val Lys Glu Tyr Lys Tyr Pro Glu Lys Leu Pro Gly Glu Leu Tyr Asp
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RAW SEQUENCE LISTING   DATE: 08/22/2005
PATENT APPLICATION: US/09/981,151D   TIME: 10:02:32

Input Set : **D:\Cura 468 SEQ list 0705.txt**
Output Set: **N:\CRF4\08222005\I981151D.raw**

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261 Ala Asn Thr Gln Cys Lys Trp Gln Phe Gly Glu Lys Ala Lys Leu Cys
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264 Met Leu Asp Phe Lys Lys Asp Ile Cys Lys Ala Leu Trp Cys His Arg
265     610              615              620
267 Ile Gly Arg Lys Cys Glu Thr Lys Phe Met Pro Ala Ala Glu Gly Thr
268 625              630              635              640
270 Ile Cys Gly His Asp Met Trp Cys Arg Gly Gly Gln Cys Val Lys Tyr
271             645              650              655
273 Gly Asp Glu Gly Pro Lys Pro Thr His Gly His Trp Ser Asp Trp Ser
274         660              665              670
276 Ser Trp Ser Pro Cys Ser Arg Thr Cys Gly Gly Gly Val Ser His Arg
277     675              680              685
279 Ser Arg Leu Cys Thr Asn Pro Asn Pro Ser His Gly Gly Lys Phe Cys
280     690              695              700
282 Glu Gly Ser Thr Arg Thr Leu Lys Leu Cys Asn Ser Gln Lys Cys Pro
283 705              710              715              720
285 Arg Asp Ser Val Asp Phe Arg Ala Ala Gln Cys Ala Glu His Asn Ser
286             725              730              735
288 Arg Arg Phe Arg Gly Arg His Tyr Lys Trp Lys Pro Gln Asp Leu Cys
289         740              745              750
291 Lys Leu Tyr Cys Ile Ala Glu Gly Phe Asp Phe Phe Phe Ser Leu Ser
292     755              760              765
294 Asn Lys Val Lys Asp Gly Thr Pro Cys Ser Glu Asp Ser Arg Asn Val
295     770              775              780
297 Cys Ile Asp Gly Ile Cys Glu Xaa Gly Cys Asp Asn Val Leu Gly Ser
298 785              790              795              800
300 Asp Ala Val Glu Asp Val Cys Gly Val Cys Asn Gly Asn Asn Ser Ala
301             805              810              815
303 Cys Thr Ile His Arg Gly Leu Tyr Thr Lys His His His Thr Asn His
304         820              825              830
306 Tyr His Met Val Thr Ile Pro Ser Gly Ala Arg Ser Ile Arg Ile Tyr
307     835              840              845
309 Glu Met Asn Val Ser Thr Ser Tyr Ile Ser Val Arg Asn Ala Leu Arg
310     850              855              860
312 Arg Tyr Tyr Leu Asn Gly His Trp Thr Val Asp Trp Pro Gly Arg Tyr
313 865              870              875              880
315 Lys Phe Ser Gly Thr Thr Phe Asp Tyr Arg Arg Ser Tyr Asn Glu Pro
316         885              890              895
318 Glu Asn Leu Ile Ala Thr Gly Pro Thr Asn Glu Thr Leu Ile Val Glu
319         900              905              910
321 Leu Leu Phe Gln Gly Arg Asn Pro Gly Val Ala Trp Glu Tyr Ser Met
322     915              920              925
324 Pro Arg Leu Gly Thr Glu Lys Gln Pro Pro Ala Gln Pro Ser Tyr Thr
325     930              935              940
327 Trp Ala Ile Val Arg Ser Glu Cys Ser Val Ser Cys Gly Gly Gly Arg
328 945              950              955              960
330 Cys Leu Pro Val Leu Leu Leu Glu Ala Ala Cys Gln Pro Ser Ala Thr
331         965              970              975
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 08/22/2005
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Input Set : **D:\Cura 468 SEQ list 0705.txt**
Output Set: **N:\CRF4\08222005\I981151D.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 857,858,2383,2983
Seq#:2; Xaa Pos. 283,792
Seq#:23; N Pos. 2196,2230,2261,2270,2295,2301
Seq#:34; Xaa Pos. 450
Seq#:65; Xaa Pos. 41

VERIFICATION SUMMARY　　　　　　DATE: 08/22/2005
PATENT APPLICATION:　**US/09/981,151D**　　TIME: 10:02:33

Input Set :　**D:\Cura 468 SEQ list 0705.txt**
Output Set:　**N:\CRF4\08222005\I981151D.raw**

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L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:840
M:341 Repeated in SeqNo=1
L:201 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:272
M:341 Repeated in SeqNo=2
L:1956 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:2160
M:341 Repeated in SeqNo=23
L:3356 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:448
L:6506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:32
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